

Figure 1

DSP-12, 1656 Base Pairs

Figure 2

DSP-12, 552 Amino Acids

MVLRLWSDTKIHLDGGFSVSTAGRMHIFKPVSQAMWSALQVLHKACEVARRHNYFP
GGVALIWATYYESCISSEQSCINEWNAMQDLESTRPDSPALFVDKPTEGERTERLIKAK
LRSIMMSQDLENVTSKEIRNELEKQMNCLKELKEFIDNEMLLILGQMDKPSLIFDHLY
LGSzWNASNLEELQGSGVITLNVIREIIONEFGIFAYHNIRVIDEETIDLLAHWNEAY
HFINKAKRNHSKCLV**HCKMGVSRS**ASTVIAYAMKEFGWPLEKAYNYVKQKRSITRPNAG
FMRQLSEYEGILDASKQRHNKLWFCQTDSSLQQPVDDPAGPGDFLPETPDGTPESQLPF
LDDAAQPGPLGPPLPCCFRLSDPLPSPEDEAGSLVHLEDPEREALLEEAAAPPAAEVHRP
ARQPQQGSGLCEKDVKKKLEFGSPKGSGSLLQVEETEREEGLGAGRWGQLPTQLDQNL
LNSENNNNSKRSCPNGMEVGRAFPAGWHTPSLPSHSNWPTSASVVGGTGTGTRHTQLIF
FYCLLWAPSSHQGPEGSFTG

Figure 3

DSP-13, 1527 Base Pairs

Figure 4

DSP-13, 509 Amino Acids

MTLSTLARKRKA
PLACTCSLGGPDMIPYFSANAVISQNA
INQLISESFLT
VKGAALFLPRGN
GS
STPRISHRRNKHAGDLQHLQAMFILLRP
EDNIRLAVRLE
STYQNRT
RYMVVV
STNGRQDTE
ES
IVLGMDFSSN
DSTCTMGLVPLW
SDTLIHLDGD
GGFSV
STDNRVHIF
KPVSVQAM
WSALQSLH
KACEV
AFLHN
FPGSL
FLW
SYV
ES
HIN
SDQ
SSV
NEWNAM
MQDV
QSH
FP
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SP
AL
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LIK
TKL
RE
IMM
MQ
KD
LEN
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IR
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LEM
QM
VC
NL
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Figure 5

A DSP13 Alternate Splice Variant, 723 Base Pairs

CTGCCC GGCTTCTAACAGGCCACTGACCGGFACTCACTGGGGACCCACGCTCTAAGTTGTTGAT
CTCTAGAACCGATTTGGAAAAGGATTTCGCTTATTGAAAGAGACAGGAFCATTCTTCTTCTT
TCCCATTTAAAGAATAATCGTTATTAAAGAATATCGTTAAAGAATAATCGTTATTCTCTCTCT
AGACCTACTGACGTGAAACGACAGAAAAGGCTAAATTAAAGAACATTAAAGGAGATC**ATGATGC**
AGAAGGATTGGAGAATATTACATCCAAAGAGATAAGAACAGAGTTGGAAATGCAAATGGTGTG
CAACTTGGGAAATTCAAGGAATTATAGACAATGAAATGATACTGATCCTTGGTCAAATGGAT
AGGCTAACAGATATTGAGCATGTGTTCTGGCTCAGAAATGGAATGCTCCAACTTAGAGG
ACTTACAGAACGGAGGGTACGGTATACTTGAATGTCAGGAGATAGATAACTTCTTCC
AGGAGTCTTGTAGTATCATAACATTGGTATATGATGAAAGAACCCATCTCTGGCGTAC
TGGAATGACAATTCAAAATTCTCTCTAAAGCAAAGAAACATGGATCTAAATGCCTTGTGCACT
GCAAAATGGGGTGAGTCGCTCAGGCTCAGCGTGAATTGCTATGCAATGAAGGAATATGGCTG
GAATCTGACGGAGCTATGACTATGTGAAAGAACAGGAAACGGTAACCAAGGCCAACCCAAAGC
TTCTGAGSACAACTGGAAAGAGTATGAGGGATCTTGCTGGCAAGCTCCTAGGCTTGATTCTG
GAGGGAGGGACAAAGGCCCTGGGGAGAGAAAAGCACAGAATTGAGTCAGTAGATCTGGTTCCAT
TCTGGTTCAUCCTCTTGCTGCAACCTGAGAAGTTACTTCACATTCTCATCCTTACCTGACC
CCATCTATAAAAT**TGAAAAT**CAAGAGATCCATCTCACAGGGTATTGTGAATAAAATGTGTTG
AATGTTATAAAAAAAAAAAAAAA

B DSP13 Alternate Splice Variant, 241 Amino Acids

MMQKDLENITSKEIRTELEMQMVCNLREFKEFIDNEMIVILGQMDSPTQIFEHVFLGSEWNASN
LEDLQNRGVRYILNVTRIEDNFFPGVFEYHNIRVYDEEATDILAYWNNTYKFISKAKKHGSKCL
VHCKMGVSRSASTVIAKAMKEYGWNLDRAYDYVKERRTVTKPNPSFMRQLEEYQGILLASFLGL
TIPDFDKFWDKKMTTFFVWVLLISIPGSSPCTDNPRHLIHTSHPVNTPSIF

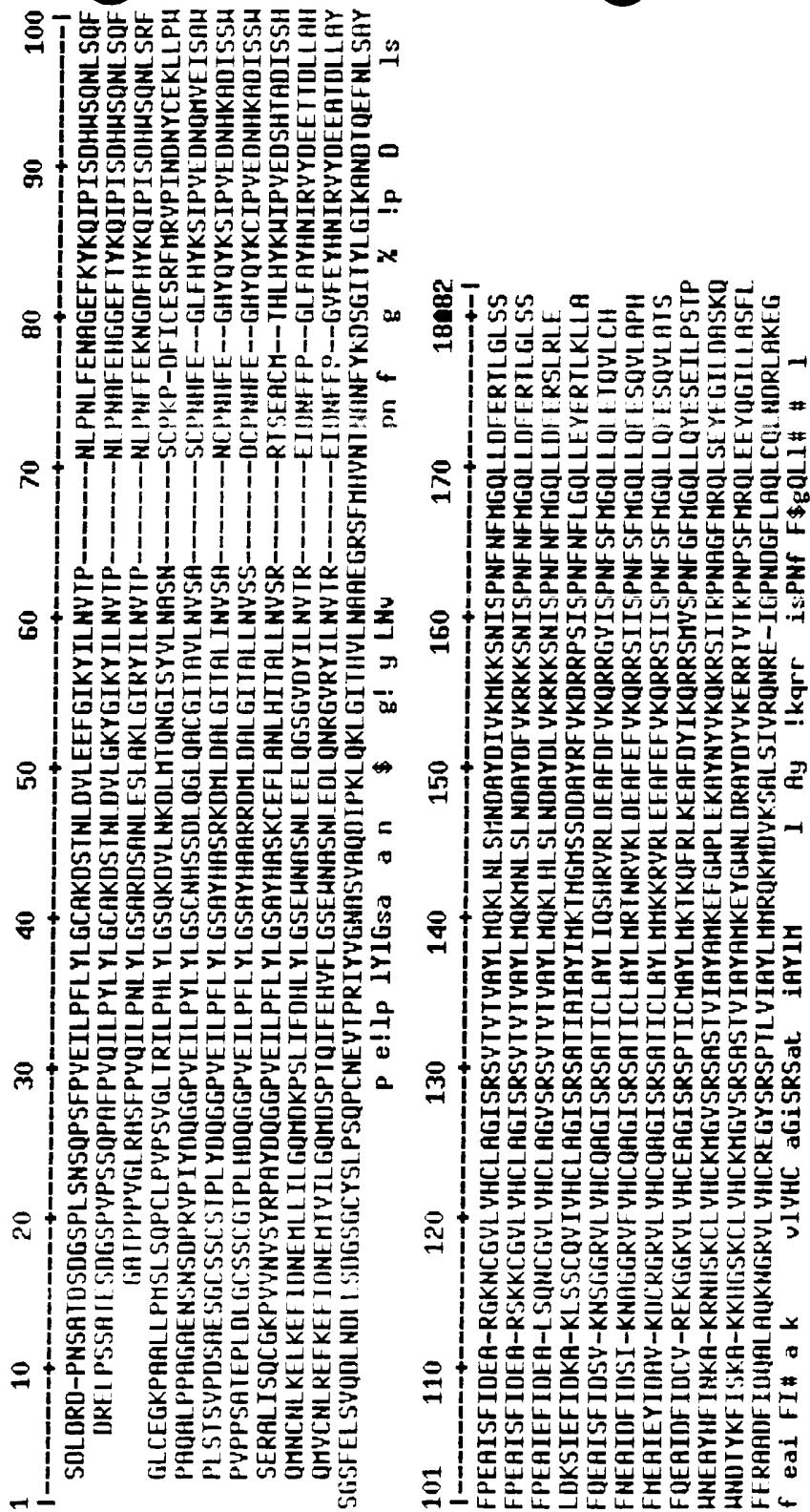


Figure 6

Alignment of DSP-12 and DSP-13

1	10	20	30	40	50	60	70	80	90	100	110
111	120	130	140	150	160	170	180	190	200	210	220
221	230	240	250	260	270	280	290	300	310	320	330
331	340	350	360	370	380	390	400	410	420	430	440
441	450	460	470	480	490	500	510	520	530	540	550
551	560	570	580	590	600	610	620	630	640	650	660
661	670	680	690	700							

Figure 7